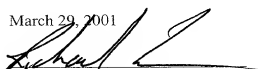


IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In the Application of:) Certification Under 37 CFR 1.10
Welcher et al.)
Serial No.: To be determined) Express Mailing Label No.
) EL 564459107 US
Filed: Herewith) I hereby certify that this paper is being
) deposited with the United States Postal
For: CD20/IgE Receptor Like) Service "EXPRESS MAIL POST
Molecules and Uses Thereof) OFFICE TO ADDRESSEE" service
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Group Art Unit: To be determined) indicated below and is addressed to the
) Commissioner for Patents, Box Patent
Examiner: To be determined) Application, Washington, D.C., 20231.
)
) March 29, 2001
)
) 
) Richard Zimmermann



STATEMENT UNDER 37 C.F.R. §1.821(f)

Box Patent Application
Commissioner for Patents
Washington, DC 20231


Sir:

I hereby state that the contents of the paper and computer readable
copies of the Sequence Listing, submitted herewith in accordance with 37 C.F.R.
§1.821(c) and (e), respectively, are the same.

Respectfully submitted,

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By


David A. Gass
Reg. No. 38,153

March 29, 2001

SEQUENCE LISTING

<110> Welcher, Andrew A.
Calzone, Frank J.

<120> CD20/IgE-Receptor Like Molecules and Uses Thereof

<130> 01017/36938A

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1 5
cac agt ccg gtg ttt ctg gta ttt cct cca gaa atc act gct tca gaa 163
His Ser Pro Val Phe Leu Val Phe Pro Pro Glu Ile Thr Ala Ser Glu
10 15 20
tat gag tcc aca gaa ctt tca gcc acg acc ttt tca act caa agc ccc 211
Tyr Glu Ser Thr Glu Leu Ser Ala Thr Thr Phe Ser Thr Gln Ser Pro
25 30 35
ttg caa aaa tta ttt gct aga aaa atg aaa atc tta ggg act atc cag 259
Leu Gln Lys Leu Phe Ala Arg Lys Met Lys Ile Leu Gly Thr Ile Gln
40 45 50
atc ctg ttt gga att atg acc ttt tct ttt gga gtt atc ttc ctt ttc 307
Ile Leu Phe Gly Ile Met Thr Phe Ser Phe Gly Val Ile Phe Leu Phe
55 60 65 70
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Thr Leu Leu Lys Pro Tyr Pro Arg Phe Pro Phe Ile Phe Leu Ser Gly
75 80 85
tat cca ttc tgg ggc tct gtt ttg ttc att aat tct gga gcc ttc cta 403
Tyr Pro Phe Trp Gly Ser Val Leu Phe Ile Asn Ser Gly Ala Phe Leu
90 95 100

att gca gtg aaa aga aaa acc aca gaa act ctg ata ata ttg agc cga 451
 Ile Ala Val Lys Arg Lys Thr Thr Glu Thr Leu Ile Ile Leu Ser Arg
 105 110 115

ata atg aat ttt ctt agt gcc ctg gga gca ata gct gga atc att ctc 499
 Ile Met Asn Phe Leu Ser Ala Leu Gly Ala Ile Ala Gly Ile Ile Leu
 120 125 130

ctc aca ttt ggt ttc atc cta gat caa aac tac att tgt ggt tat tct 547
 Leu Thr Phe Gly Phe Ile Leu Asp Gln Asn Tyr Ile Cys Gly Tyr Ser
 135 140 145 150

cac caa aat agt cag tgt aag gct gtt act gtc ctg ttc ttg gga att 595
 His Gln Asn Ser Gln Cys Lys Ala Val Thr Val Leu Phe Leu Gly Ile
 155 160 165

ttg att aca ttg atg act ttc agc att att gaa tta ttc att tct ctg 643
 Leu Ile Thr Leu Met Thr Phe Ser Ile Ile Glu Leu Phe Ile Ser Leu
 170 175 180

cct ttc tca att ttg ggg tgc cac tca gag gat tgt gat tgt gaa caa 691
 Pro Phe Ser Ile Leu Gly Cys His Ser Glu Asp Cys Asp Cys Glu Gln
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 35 40 45

Ile Leu Gly Thr Ile Gln Ile Leu Phe Gly Ile Met Thr Phe Ser Phe
 50 55 60

Gly Val Ile Phe Leu Phe Thr Leu Leu Lys Pro Tyr Pro Arg Phe Pro
 65 70 75 80

Phe Ile Phe Leu Ser Gly Tyr Pro Phe Trp Gly Ser Val Leu Phe Ile
 85 90 95

Asn Ser Gly Ala Phe Leu Ile Ala Val Lys Arg Lys Thr Thr Glu Thr
 100 105 110

Leu Ile Ile Leu Ser Arg Ile Met Asn Phe Leu Ser Ala Leu Gly Ala
 115 120 125

Ile Ala Gly Ile Ile Leu Leu Thr Phe Gly Phe Ile Leu Asp Gln Asn
 130 135 140

Tyr Ile Cys Gly Tyr Ser His Gln Asn Ser Gln Cys Lys Ala Val Thr
 145 150 155 160
 Val Leu Phe Leu Gly Ile Leu Ile Thr Leu Met Thr Phe Ser Ile Ile
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 Met Leu Leu
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 caa tcc caa acc atg ggg gtt tct cac agc ttt aca cca aag ggc atc 163
 Gln Ser Gln Thr Met Gly Val Ser His Ser Phe Thr Pro Lys Gly Ile
 5 10 15
 act atc cct caa aga gag aaa cct gga cac atg tac caa aac gaa gat 211
 Thr Ile Pro Gln Arg Glu Lys Pro Gly His Met Tyr Gln Asn Glu Asp
 20 25 30 35
 tac ctg cag aac ggg ctg cca aca gaa acc acc gtt ctt ggg act gtc 259
 Tyr Leu Gln Asn Gly Leu Pro Thr Glu Thr Thr Val Leu Gly Thr Val
 40 45 50
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 Gln Ile Leu Cys Cys Leu Leu Ile Ser Ser Leu Gly Ala Ile Leu Val
 55 60 65
 ttt gct ccc tac ccc tcc cac ttc aat cca gca att tcc acc act ttg 355
 Phe Ala Pro Tyr Pro Ser His Phe Asn Pro Ala Ile Ser Thr Thr Leu
 70 75 80
 atg tct ggg tac cca ttt tta gga gct ctg tgt ttt ggc att act gga 403
 Met Ser Gly Tyr Pro Phe Leu Gly Ala Leu Cys Phe Gly Ile Thr Gly
 85 90 95
 tcc ctc tca att atc tct gga aaa caa tca act aag ccc ttt gac ctg 451
 Ser Leu Ser Ile Ile Ser Gly Lys Gln Ser Thr Lys Pro Phe Asp Leu
 100 105 110 115
 agc agc ttg acc tca aat gca gtg agt tct gtt act gca gga gca ggc 499
 Ser Ser Leu Thr Ser Asn Ala Val Ser Ser Val Thr Ala Gly Ala Gly
 120 125 130

ctc ttc ctc ctt gct gac agc atg gta gcc ctg agg act gcc tct caa 547
Leu Phe Leu Leu Ala Asp Ser Met Val Ala Leu Arg Thr Ala Ser Gln
135 140 145

cat tgt ggc tca gaa atg gat tat cta tcc tca ttg cct tat tcg gag 595
His Cys Gly Ser Glu Met Asp Tyr Leu Ser Ser Leu Pro Tyr Ser Glu
150 155 160

tac tat tat cca ata tat gaa atc aaa gat tgt ctc ctg acc agt gtc 643
Tyr Tyr Tyr Pro Ile Tyr Glu Ile Lys Asp Cys Leu Leu Thr Ser Val
165 170 175

agt tta aca ggt gtc cta gtg gtg atg ctc atc ttc act gtg ctg gag 691
Ser Leu Thr Gly Val Leu Val Val Met Leu Ile Phe Thr Val Leu Glu
180 185 190 195

ctc tta tta gct gca tac agt tct gtc ttt tgg tgg aaa cag ctc tac 739
Leu Leu Leu Ala Tyr Ser Ser Val Phe Trp Trp Lys Gln Leu Tyr
200 205 210

tcc aac aac cct ggg agt tca ttt tcc tcg acc cag tca caa gat cat 787
Ser Asn Asn Pro Gly Ser Ser Phe Ser Ser Thr Gln Ser Gln Asp His
215 220 225

atc caa cag gtc aaa aag agt tct tca cgg tct tgg ata taagtaactc 836
Ile Gln Gln Val Lys Lys Ser Ser Ser Arg Ser Trp Ile
230 235 240

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Asn Glu Asp Tyr Leu Gln Asn Gly Leu Pro Thr Glu Thr Thr Val Leu
35 40 45

Gly Thr Val Gln Ile Leu Cys Cys Leu Leu Ile Ser Ser Leu Gly Ala
50 55 60

Ile Leu Val Phe Ala Pro Tyr Pro Ser His Phe Asn Pro Ala Ile Ser
65 70 75 80

Thr Thr Leu Met Ser Gly Tyr Pro Phe Leu Gly Ala Leu Cys Phe Gly
85 90 95

Ile Thr Gly Ser Leu Ser Ile Ile Ser Gly Lys Gln Ser Thr Lys Pro
100 105 110

Phe Asp Leu Ser Ser Leu Thr Ser Asn Ala Val Ser Ser Val Thr Ala
115 120 125

Gly Ala Gly Leu Phe Leu Leu Ala Asp Ser Met Val Ala Leu Arg Thr
130 135 140

Ala Ser Gln His Cys Gly Ser Glu Met Asp Tyr Leu Ser Ser Leu Pro
145 150 155 160

Tyr Ser Glu Tyr Tyr Tyr Pro Ile Tyr Glu Ile Lys Asp Cys Leu Leu
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Thr Ser Val Ser Leu Thr Gly Val Leu Val Val Met Leu Ile Phe Thr
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Val Leu Glu Leu Leu Leu Ala Ala Tyr Ser Ser Val Phe Trp Trp Lys
195 200 205

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<211> 23

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<210> 16
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Arg Arg Arg